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Supporting information for article:

Crystal structures of *Escherichia coli* branching enzyme in complex with cyclodextrins

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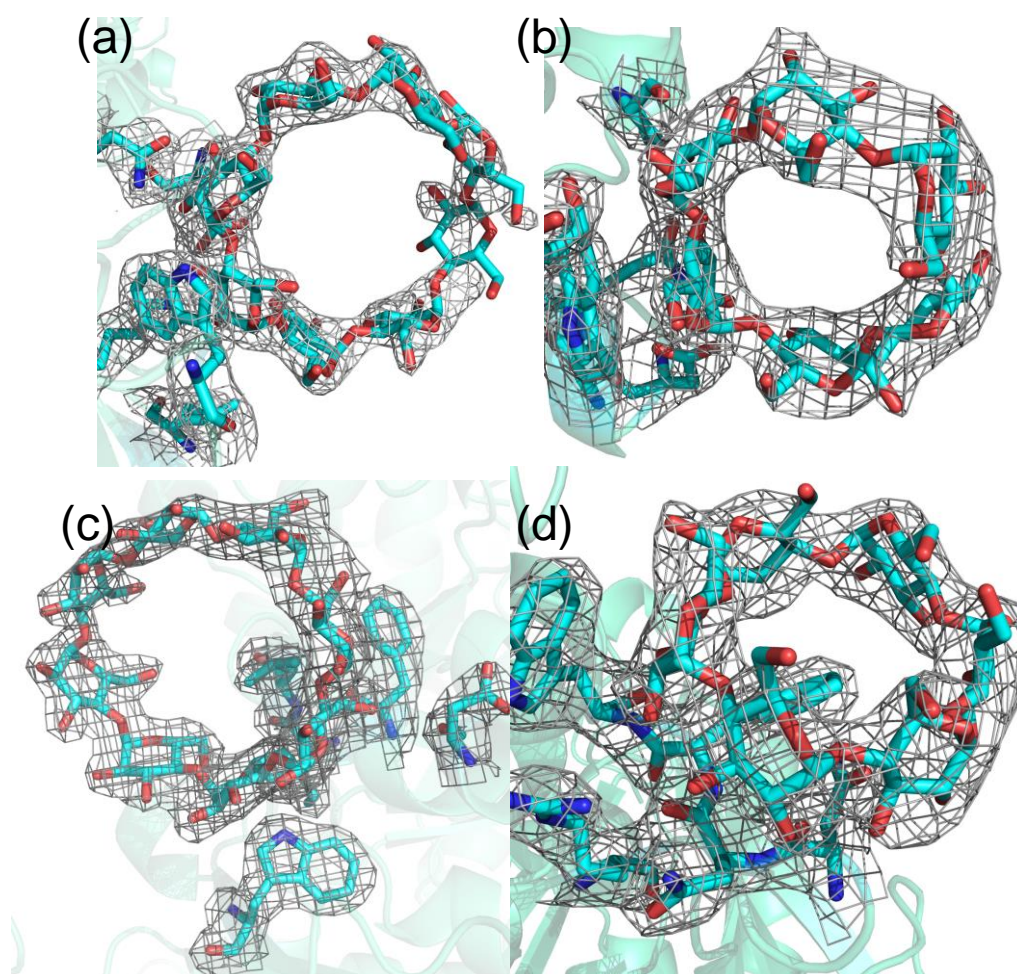


Figure S1 Electron density maps contoured around each of the four CD binding sites. (a) Electron density, contoured at 1σ , of γ -CD bound in site IV (molecule D). (b) Electron density, contoured at 1σ of α -CD bound in site V (molecule D). (c) Electron density, contoured at 1σ of γ -CD bound in site VI (molecule C). (d)